



The Future of Forensic DNA Analysis

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Presentation & Written Article Outline

- **Introduction to NIST and recent U.S. activities**
 - National Commission on Forensic Science (NCFS)
 - Organization of Scientific Area Committees (OSAC)
- **DNA capabilities**
- **The Past Reviewed**
 - Major themes and time periods
 - Research leadership
- **The Present Considered**
 - Genetic marker systems
 - DNA database growth and use in the U.S.
 - Critical challenges faced today
- **The Future Predicted**
 - *Faster* results
 - *Higher* sensitivity and information content
 - *Stronger* conclusions with challenging samples

Background Information on NIST

- Started in 1901 with roots back to the Constitution
- Name changed to **National Institute of Standards and Technology (NIST)** from National Bureau of Standards in 1988
- Primary campus in Gaithersburg, Maryland (just outside of Washington, D.C.)
- Part of the U.S. Department of Commerce
- >3,000 employees and >2,000 associates
- Supply >1300 reference materials
- Defines official time for the U.S.



Types of Standards

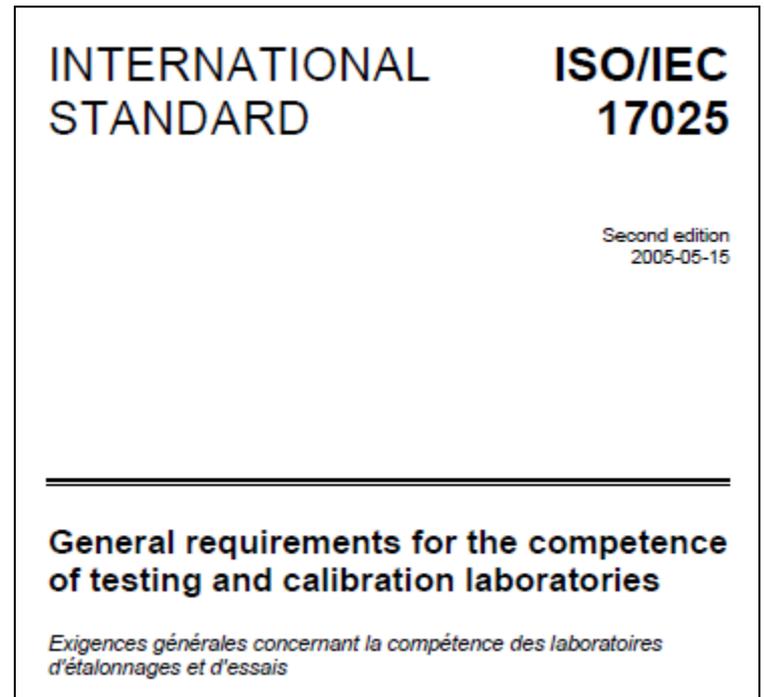
physical (measurement) standards



Certified reference material to aid with calibration of measurements

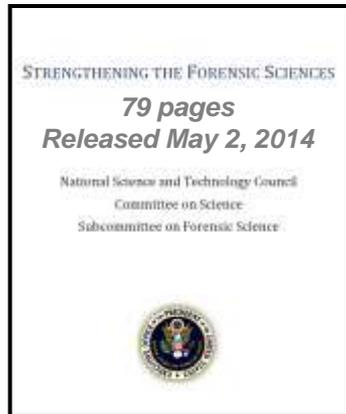
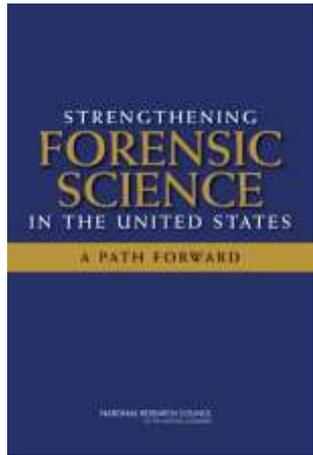
<http://www.nist.gov/srm/>

documentary (technical) standards



Specific requirements for the operation of a laboratory related to management system and competence

NCFS and OSAC: U.S. Efforts to Strengthen Forensic Science



http://www.whitehouse.gov/sites/default/files/microsites/ostp/NSTC/strengthening_the_forensic_sciences_may_-_2014.pdf

- National Academy of Sciences (**NAS**) **report** issued in Feb 2009
- White House **Subcommittee on Forensic Science** (SoFS) operated from July 2009 to Dec 2012

DOJ/NIST Partnership (announced Feb 2013)

1. **NCFS** (National Commission on Forensic Science)
 - First meeting held February 3-4, 2014 in Washington DC
2. **OSAC** (Organization of Scientific Area Committees)
 - **Being organized**; first public meetings to be held Feb 2015

National Commission on Forensic Science (NCFS)

www.justice.gov/ncfs

Policy-focused

NCFS Leadership

National Commission on Forensic Science Home

Members
Meetings
Subcommittees
Press Room
Contact the Commission



In 2013, the Department of Justice (DOJ) established the National Commission on Forensic Science, in partnership with the National Institute of Standards and Technology (NIST), to enhance the practice and improve the reliability of forensic science. This unique partnership draws upon each agency's core strengths to promote scientific validity, reduce fragmentation, and improve federal coordination of forensic science.

The Commission is co-chaired by Deputy Attorney General James M. Cole and NIST Acting Director and Acting Under Secretary of Commerce for Standards and Technology, Willie E. May. Nelson Santos, Deputy Assistant Administrator for the Office of Forensic Sciences at the Drug Enforcement Administration, and John M. Butler, Special Assistant to the NIST Director for forensic science, serve as vice-chairs. Brette Steele, Senior Advisor on Forensic Science and Senior Counsel to the Deputy Attorney General serves as the Designated Federal Officer and Robin Jones, Consultant within the Department of Justice, serves as Program Manager.

The Commission includes federal, state and local forensic science service providers; research scientists and academics; law enforcement officials; prosecutors, defense attorneys and judges; and other stakeholders from across the country.

GENERAL INFORMATION NATIONAL COMMISSION ON FORENSIC SCIENCE

CONTACT

Brette Steele
Brette.L.Steele@usdoj.gov

By Phone:
(202) 305-0180

31 voting and 8 *ex-officio* members

Last meeting (5th): January 29-30, 2015

Next meeting (6th): April 30-May 1, 2015



Sally Q. Yates
Acting
Deputy Attorney General
DOJ Co-Chair



Willie E. May
Acting
Director of NIST
NIST Co-Chair



Nelson A. Santos
Vice-Chair (DOJ)



John M. Butler
Vice-Chair (NIST)

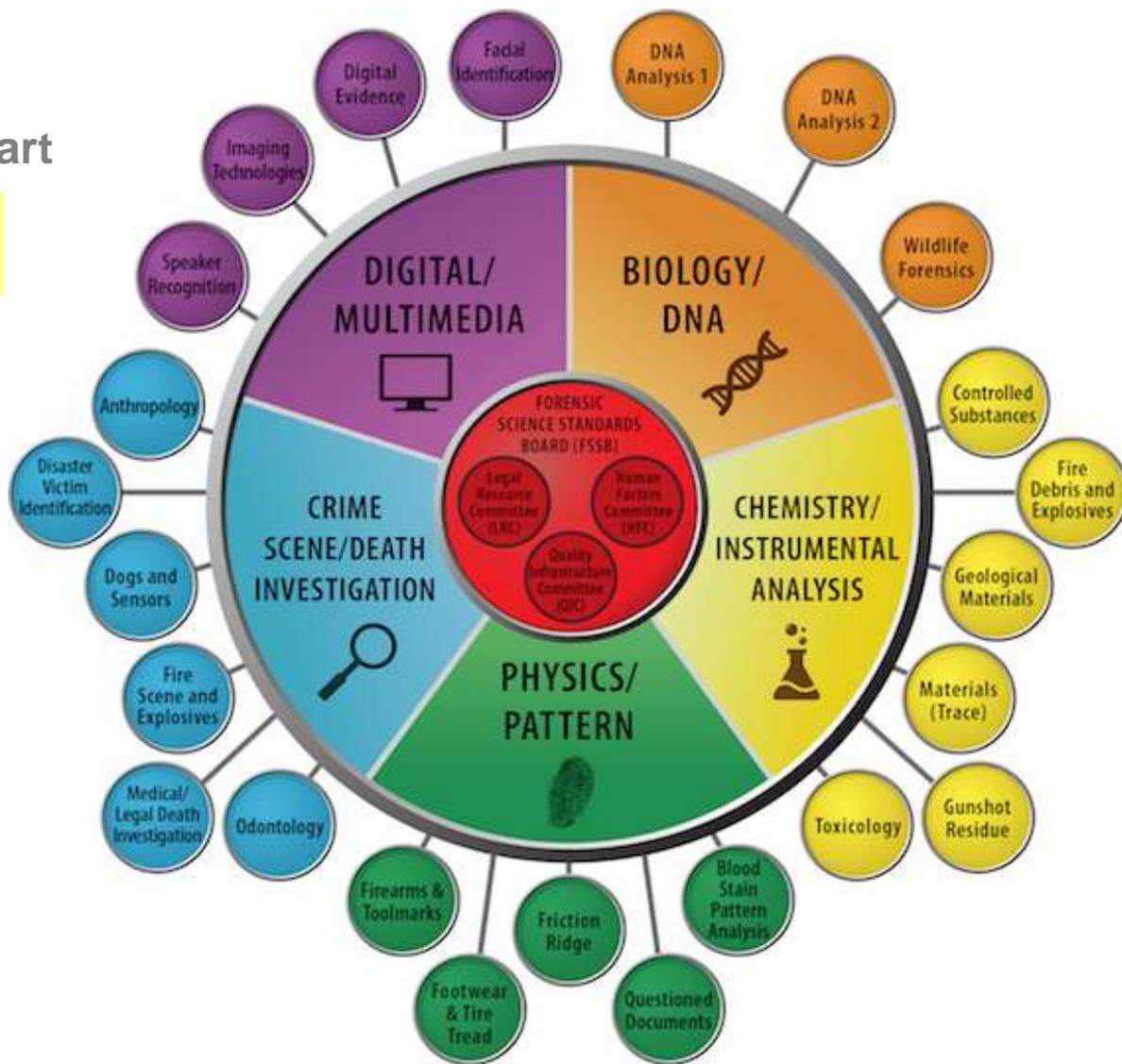
Organization of Scientific Area Committees

OSAC

Functional Organization Chart

Practice-focused

542 members and >1000 affiliates as subject matter experts participating in 24 subcommittees, 5 scientific areas, 3 resource committees (legal, quality, human factors), and 1 governing board (Forensic Science Standards Board)



<http://www.nist.gov/forensics/osac/index.cfm>

Initial membership
finalized Dec 22, 2014

Overview of Standards involved in the Forensic Science Enterprise

ILAC-G19:08/2014

Modules in a Forensic Science Process

ISO/IEC 17024

Certification
Provider

ISO/IEC 17043

Proficiency Test
Provider

ISO/IEC 17011

Accrediting Body

Forensic Testing Laboratory

Technical
standard

ISO/IEC 17025

Supplemental
material

DNA

Latents

Firearms

...

QAS

Police Agency

ISO/IEC 17020

Crime scene unit

M.E./Coroner

ISO 15189

Autopsy

Virtopsy

OSAC work will help provide supplemental materials and new technical standards

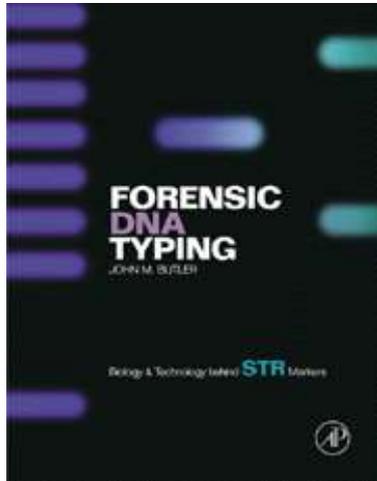
Accreditation to appropriate quality standards should provide confidence for all stakeholders in what is being done

Forensic DNA Typing Textbooks Have Set the Standard for the Field

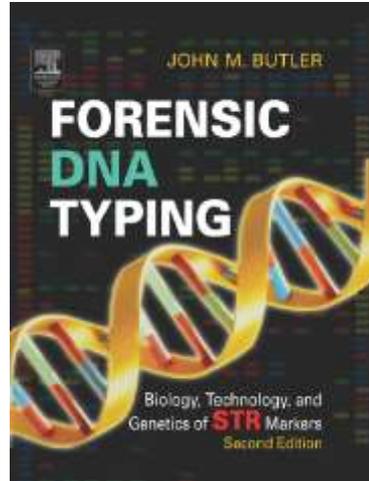
1st Edition

2nd Edition

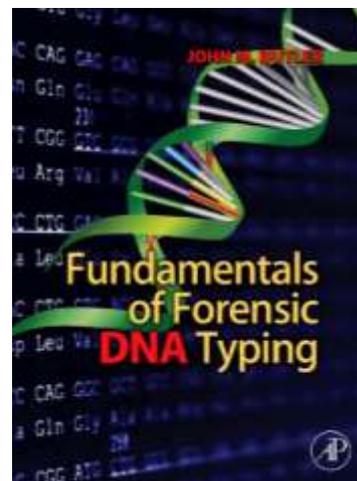
3rd Edition (3 volumes)



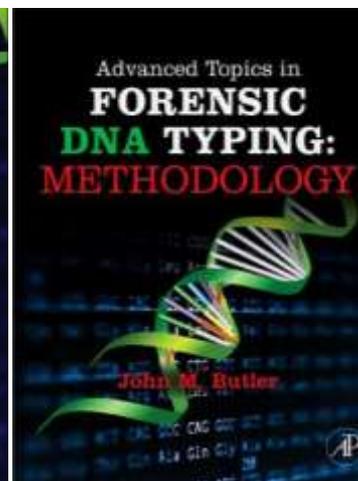
Jan 2001
335 pages



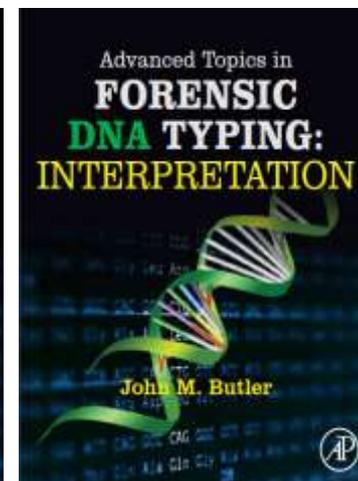
Feb 2005
688 pages



Sept 2009
520 pages



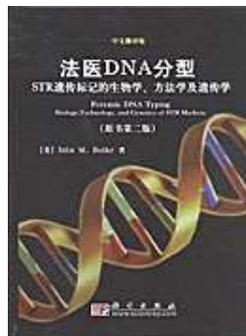
Aug 2011
704 pages



Oct 2014
604 pages

Language Editions

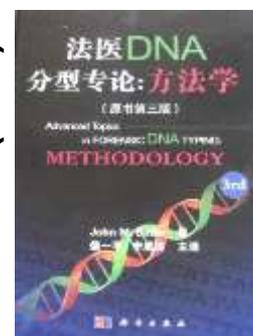
Chinese (2007)



Japanese (2009)



Chinese (2013)



Steps in Forensic DNA Analysis

Gathering the Data

Understanding Results Obtained & Sharing Them

Collection/Storage/
Characterization

Extraction/
Quantitation

Amplification/
Marker Sets

Separation/
Detection

Data

Stats

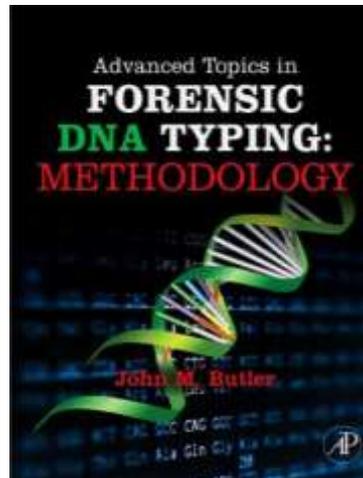
Report

Interpretation

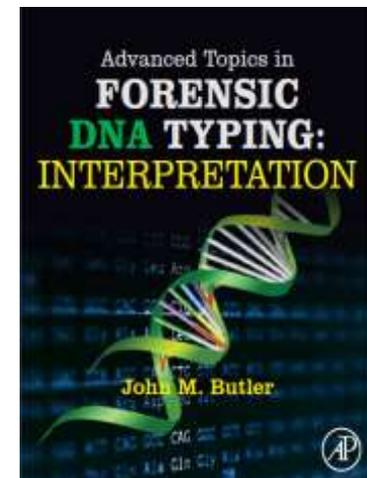
Advanced Topics: Methodology

Advanced Topics: Interpretation

>1300 pages of information with >5000 references cited in these two books



August 2011



October 2014

DNA Capabilities to Aid Forensic Investigations

1. The **ability to identify the perpetrator**
2. Weight-of-evidence based on established genetic principles and statistics (Hardy-Weinberg 1908)
3. Established characteristics of genetic inheritance enables close **biological relatives** to be used for reference points using kinship associations
4. Superb **sensitivity** with PCR amplification (opens the possibility for contamination)
5. Well-established **quality assurance measures**
6. New **technology development** aided by genomics

Successful interpretation of DNA (Q-to-K comparison) depends on quality of the crime scene evidence (Q) and availability of suitable reference samples (K)

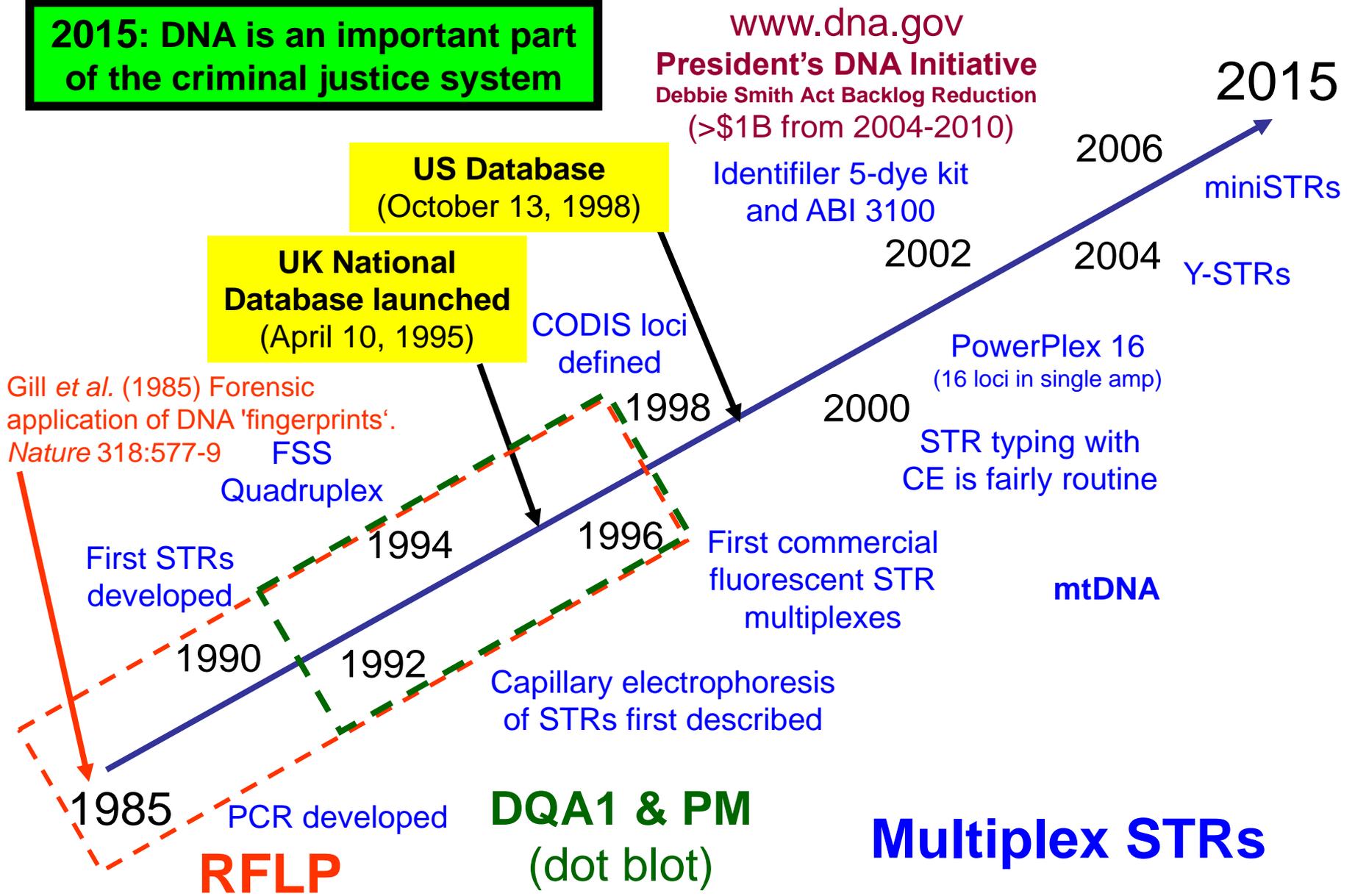
The
Past
Reviewed

Stages of Forensic DNA Progression

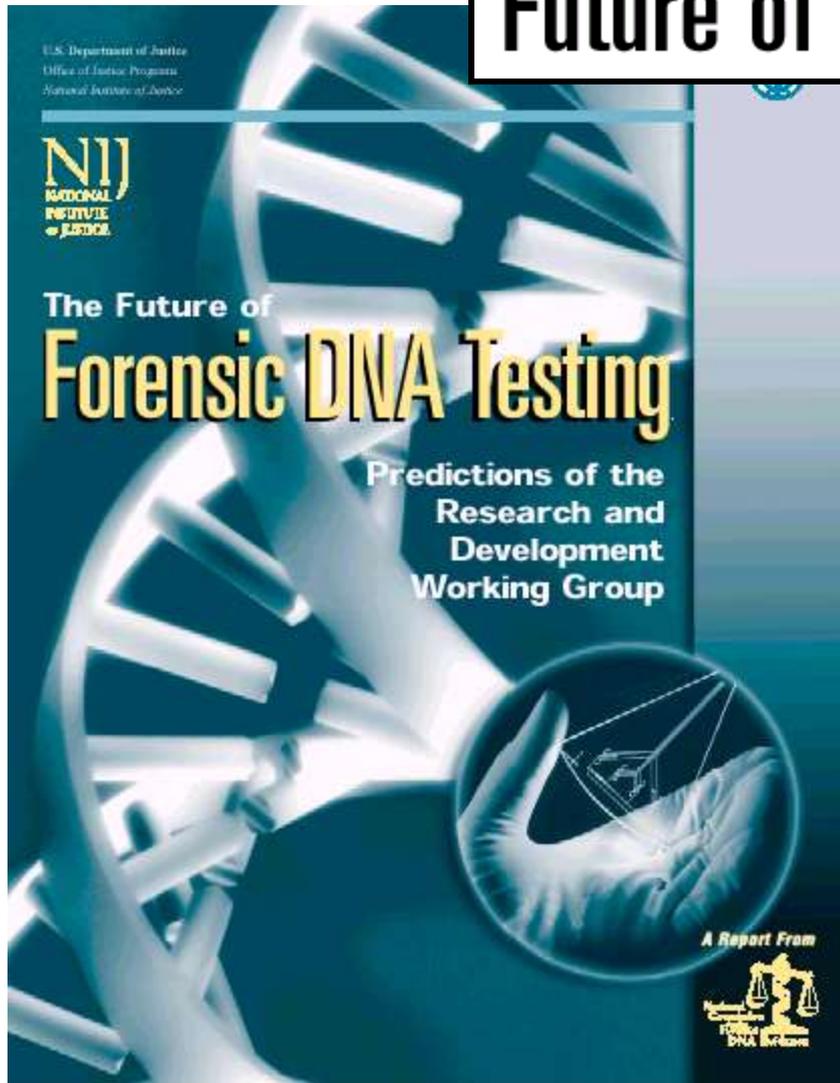
Stages	Time Frame	Description
Exploration	1985 - 1995	Beginnings, different methods tried (RFLP and early PCR)
Stabilization	1995 - 2005	Standardization to STRs, selection of core loci, implementation of Quality Assurance Standards
Growth	2005 - 2015	Rapid growth of DNA databases, extended applications pursued
<i>Sophistication</i>	<i>2015 to 2025 and beyond</i>	<i>Expanding tools available, confronting privacy concerns</i>

History of Forensic DNA Testing

2015: DNA is an important part of the criminal justice system



National Commission on the Future of DNA Evidence



- Report published in Nov 2000
- Asked to estimate where DNA testing would be 2, 5, and 10 years into the future

Conclusions

STR typing is here to stay for a few years because of DNA databases that have grown to contain millions of profiles

Research Leadership in Forensic DNA

- During its existence, the **UK Forensic Science Service** played an important role in the development and application of forensic DNA techniques
- Other important centers of research include:
 - University of Innsbruck (Austria)
 - University of Copenhagen (Denmark)
 - University of Santiago de Compostella (Spain)
 - NIST Applied Genetics Group (USA)
 - University of North Texas Health Science Center (USA)
 - Netherlands Forensic Institute (Holland)
 - Institute of Environmental Science and Research (New Zealand)

The
Present
Considered

Current U.S. National DNA Database

As of December 2014, over **13 million samples**

- 11,548,720 offender DNA profiles
- 1,303,454 arrestee DNA profiles
- 601,664 forensic profiles

As of October 2014, China had >420 labs and >25 million STR profiles in their DNA database

- Has produced **270,326 hits** to help solve cases
- <http://www.fbi.gov/about-us/lab/biometric-analysis/codis/ndis-statistics>

Growth of DNA Databases

- Expanded laws now enable more offenders to be included (32 states and federal government have laws to collect from arrestees)
 - Has contributed to sample backlogs
- Have benefited from significant federal funding since 2004 (>\$1 billion for backlog reduction)
- **Have effectively locked technology with core STR markers used** to generate DNA profiles that now number in the millions

U.S. Supreme Court
decision (June 2013)
in *Maryland v King*

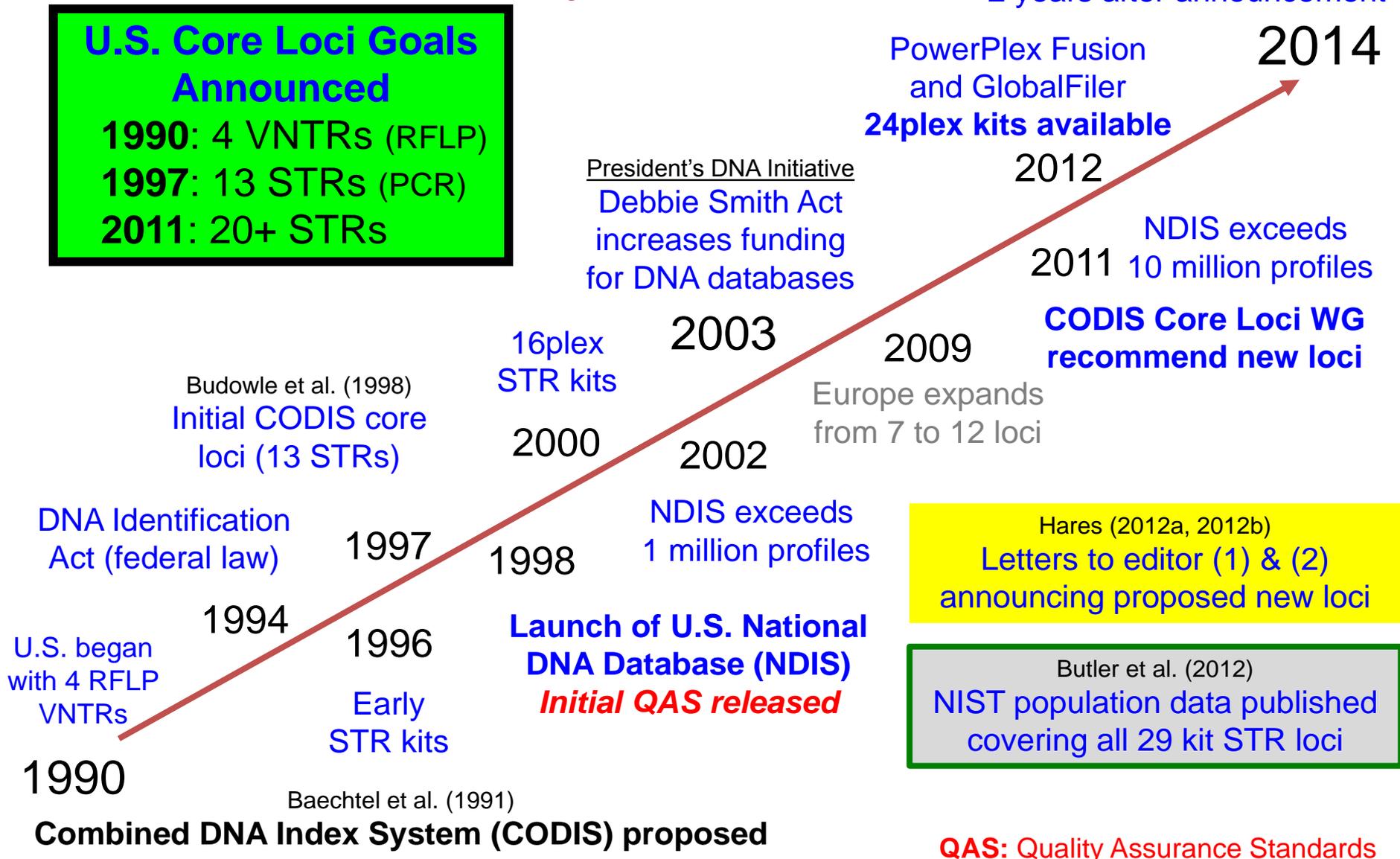
U.S. Core Loci Expansion Efforts

More loci added as databases grew...

Implementation to be required
2 years after announcement

U.S. Core Loci Goals Announced

1990: 4 VNTRs (RFLP)
1997: 13 STRs (PCR)
2011: 20+ STRs



QAS: Quality Assurance Standards

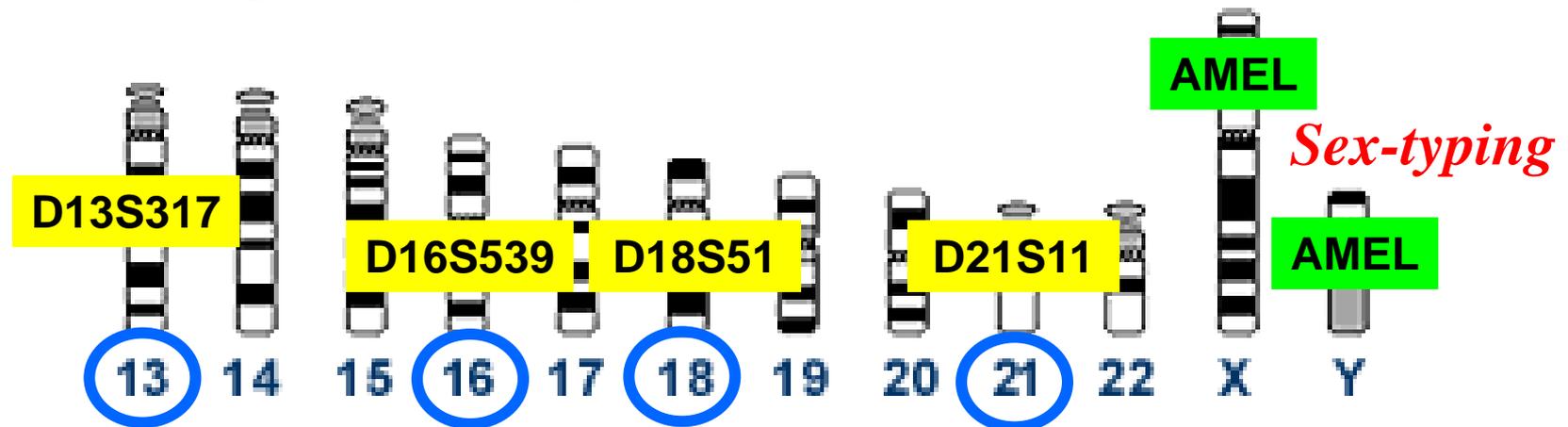
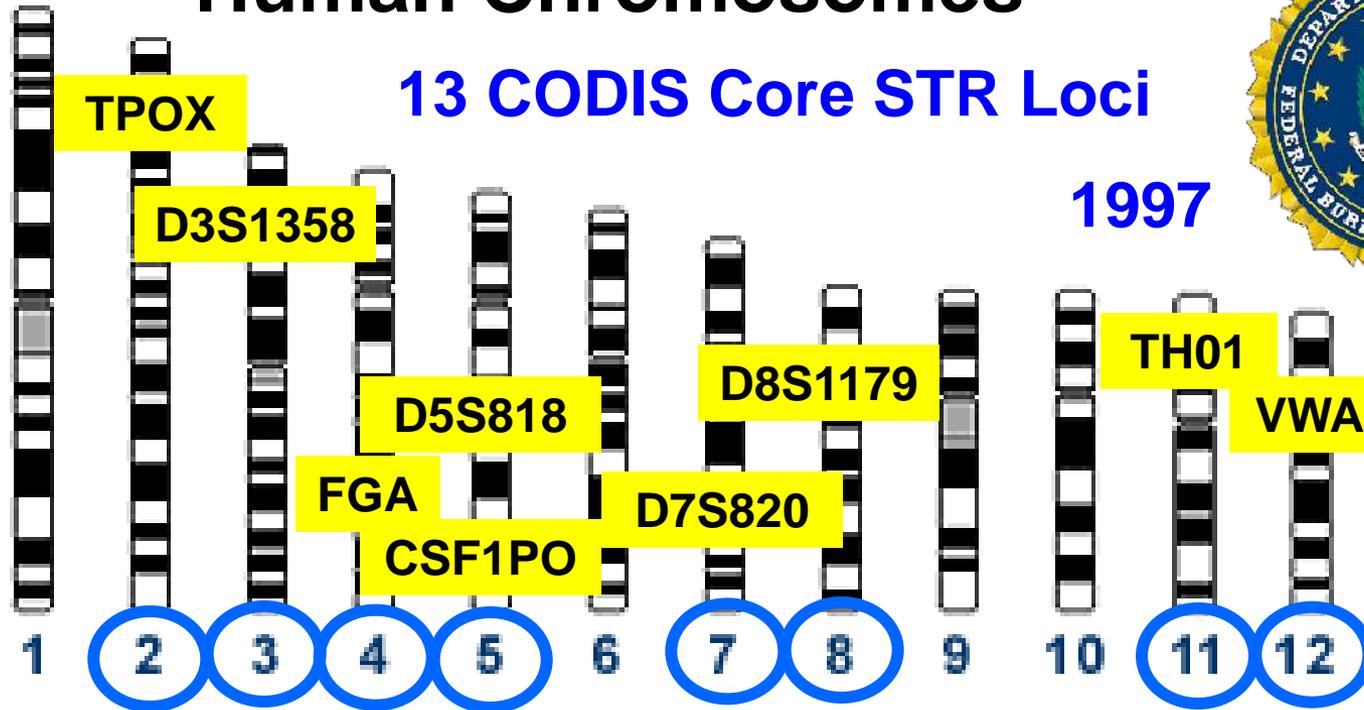
Position of Forensic STR Markers on Human Chromosomes

Core STR Loci for the United States



13 CODIS Core STR Loci

1997



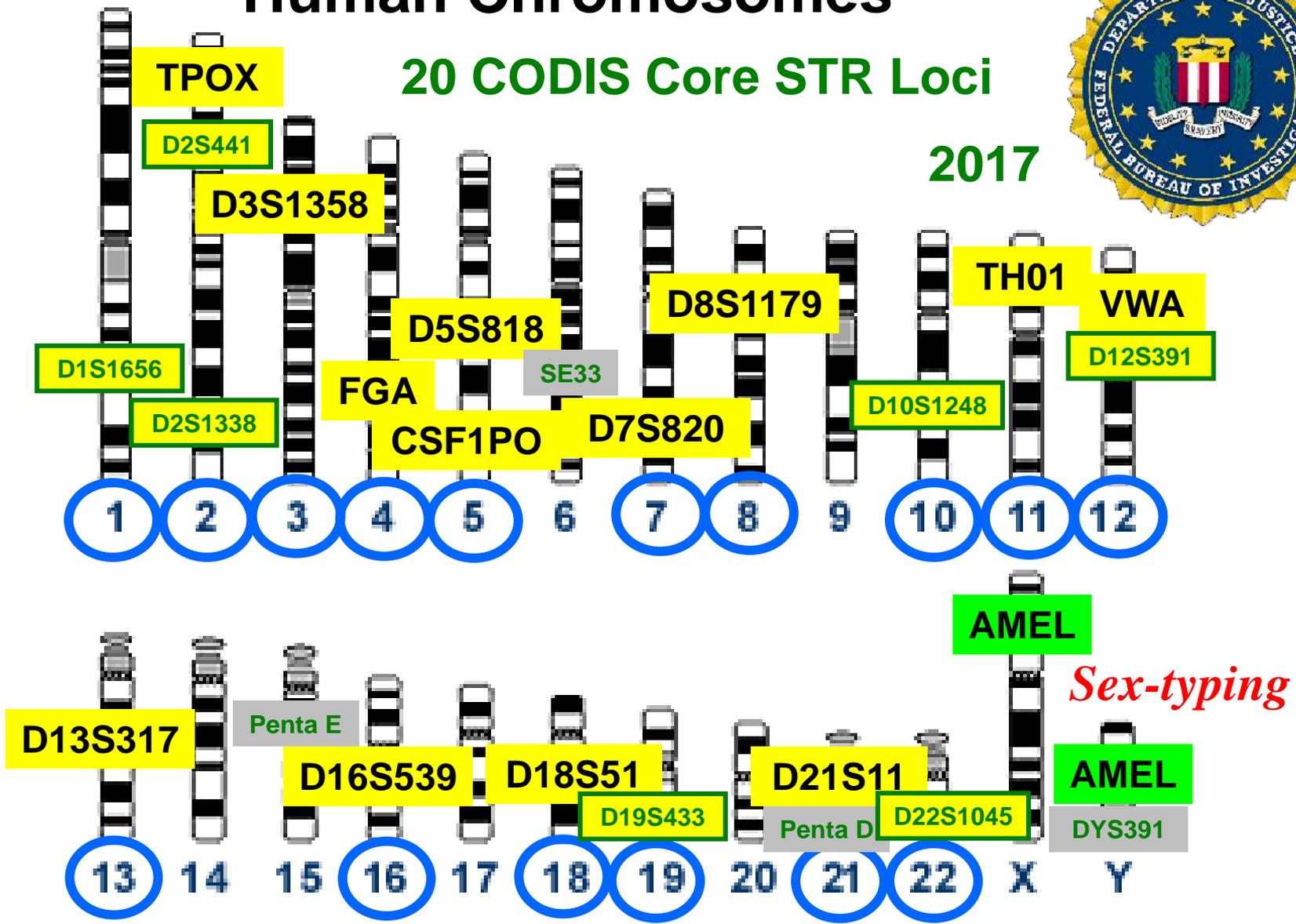
Expanded Core STR Loci for the United States

Position of Forensic STR Markers on Human Chromosomes



20 CODIS Core STR Loci

2017



Relative Sizes of STR Loci in 24plex Kits

100 bp

200 bp

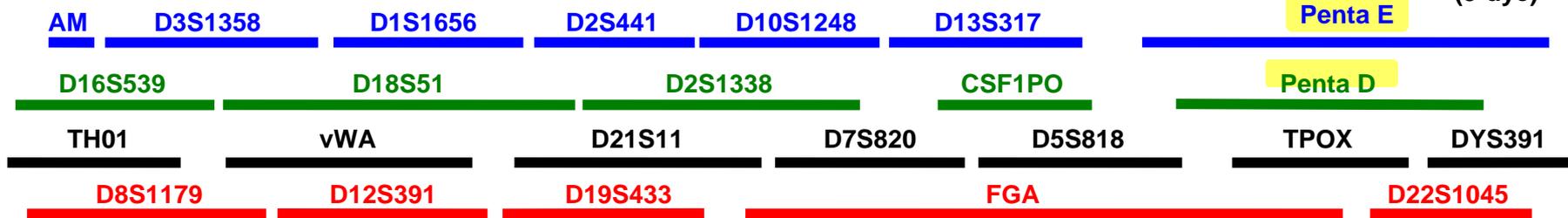
300 bp

400 bp

24plex
(5-dye)

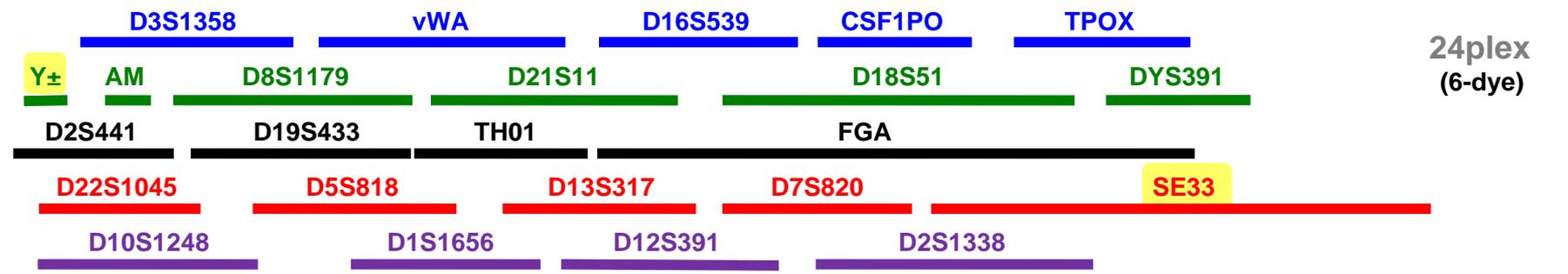
2012

PowerPlex Fusion



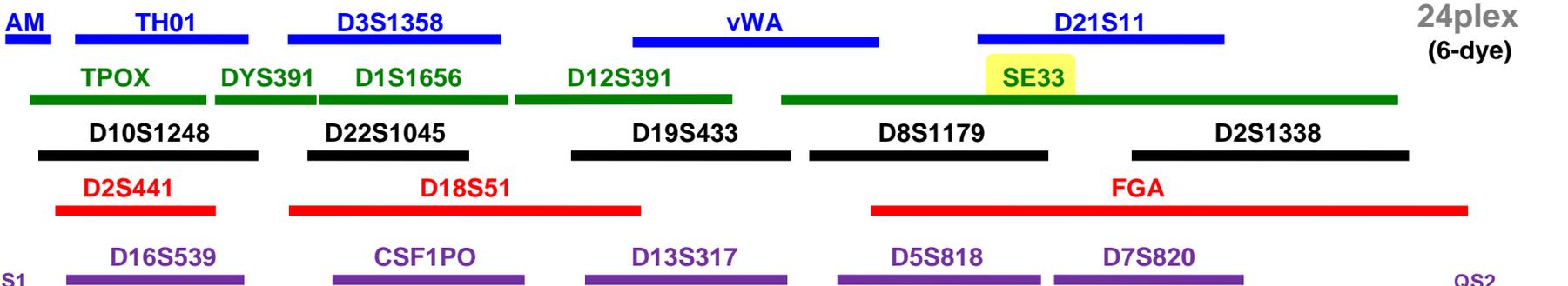
2013

GlobalFiler



2015

Investigator 24plex



The
Future
Predicted

New Trends in Forensic DNA

- ***Faster results:*** Rapid DNA capabilities and new sample-to-answer integrated instruments
- ***Higher sensitivity:*** New assays lowering the limits of detection, which makes interpretation more challenging
- ***Higher information content:*** Next-generation sequencing (NGS) for more markers & STR allele information
- ***Stronger conclusions:*** Mixture interpretation with probabilistic genotyping models

Rapid DNA Efforts



Pete Vallone Erica Butts

Accelerated Nuclear DNA Equipment (ANDE) developed by **NetBio**



<http://ishinews.com/wp-content/uploads/2012/10/Rapid-DNA-Miles-1.58MB.pdf>

RapidHIT 200 developed by **IntegenX**

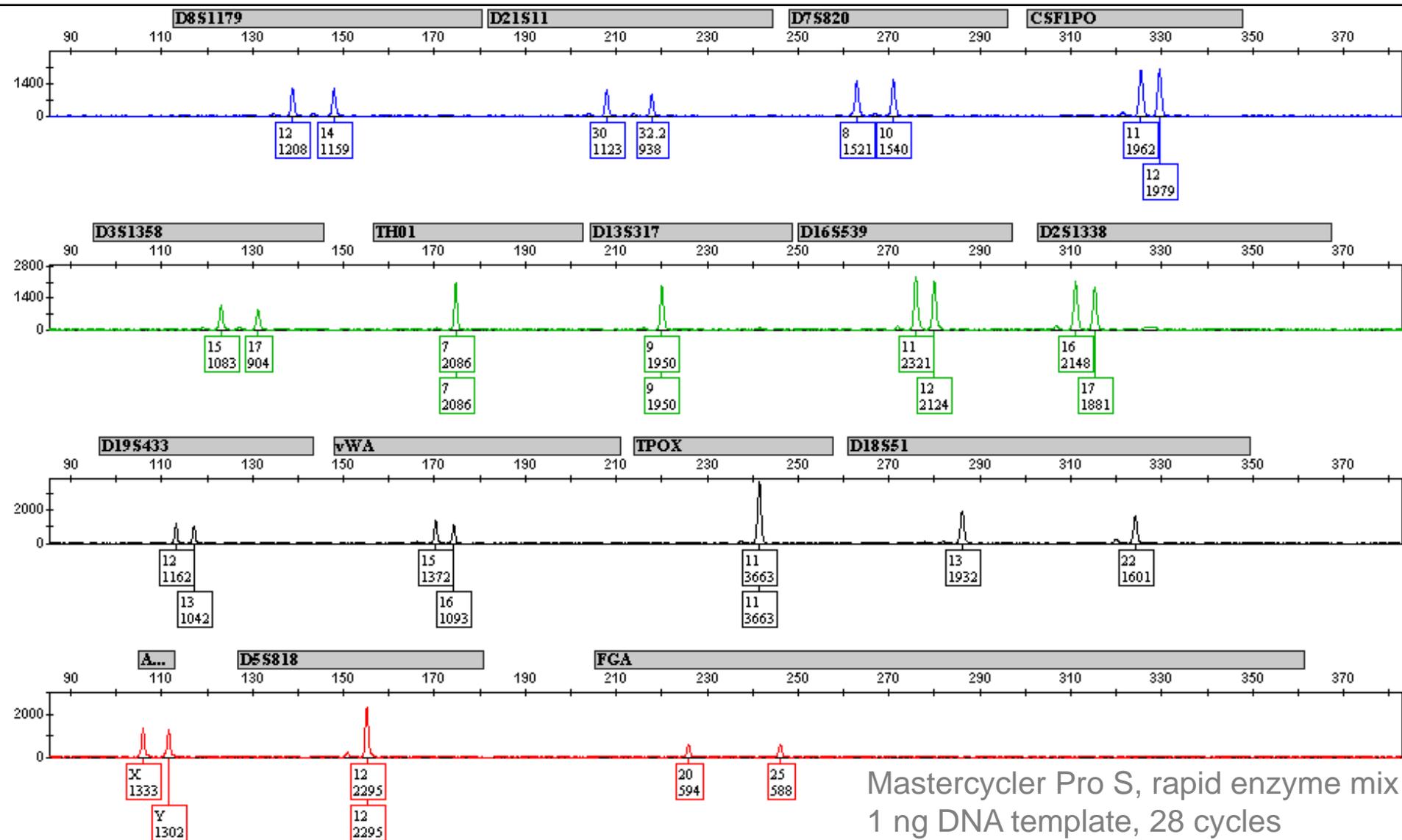


<http://integenx.com/wp-content/uploads/2010/06/RapidHIT-200.png>

- Evaluating ANDE (NetBio) and IntegenX rapid DNA instruments
 - both instruments are capable of swab in → STR profile out in less than 90 minutes without user intervention
- Exploring rapid DNA techniques including direct PCR and rapid PCR
 - STR profiles generated in <2 hours with standard lab equipment and rapid protocols
 - See ISHI 2012 poster available on STRBase “Rapid DNA Testing Approaches for Reference Samples”

Fastest results swab-to-profile (Identifiler): 57 minutes

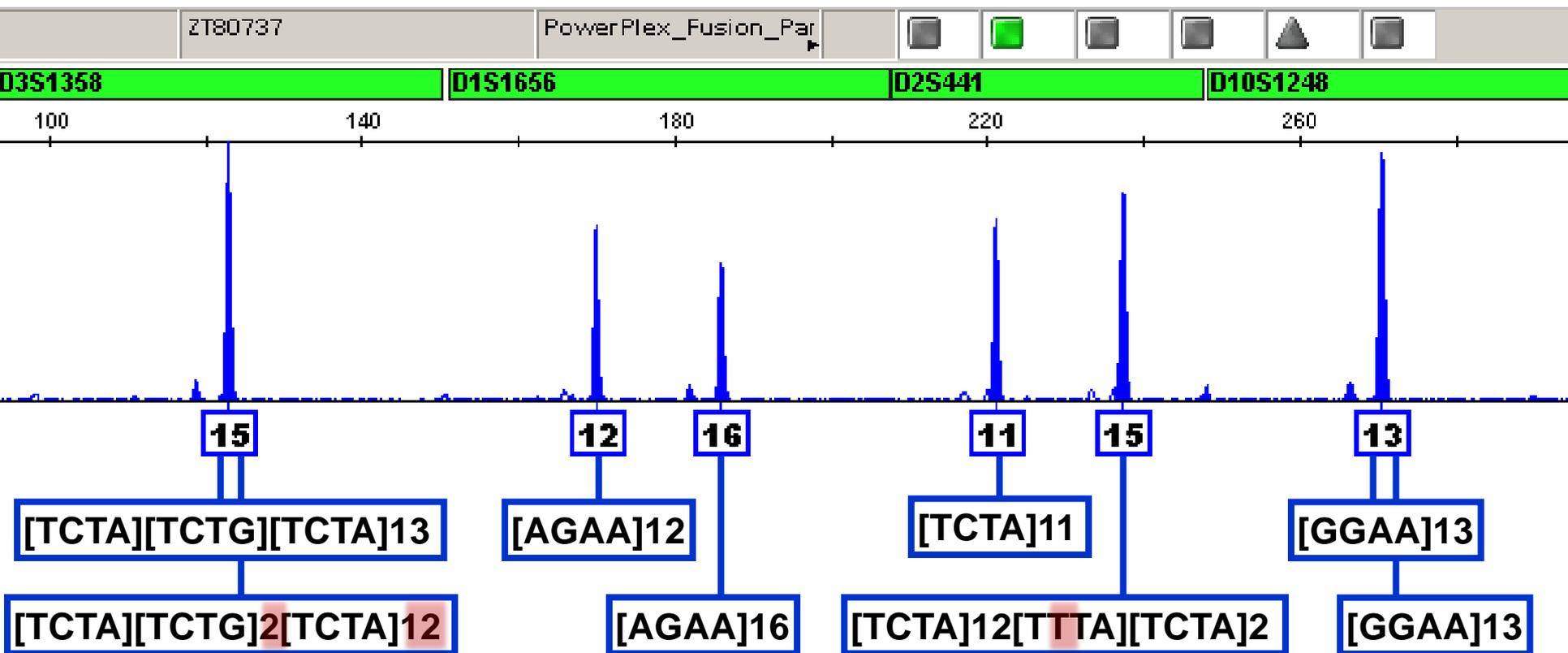
Full Identifiler STR Profile with 19 min PCR



Mastercycler Pro S, rapid enzyme mix
1 ng DNA template, 28 cycles

Next Generation Sequencing (NGS)

- **Higher information content** with sequence data
 - Expanded number of STR loci and other genetic markers such as SNPs and InDels
 - New markers may enable additional applications (e.g., biogeographical ancestry and phenotypic prediction)
 - **Deeper depth of information on STR alleles**
 - For example, eight different sequence versions of D12S391 alleles among 197 samples examined (Gelardi et al. 2014)
- **Significant challenges with BIG data**
 - STR allele nomenclature issues
 - Data storage (do you retain terabytes of data?)
 - Data analysis time will increase...
 - Privacy concerns with additional genomic information



Sequence-Based Heterozygote: A locus that appears homozygous in length-based measurements (such as CE), but is heterozygous by sequence

CSI:

Compromised Sample Improvements

- Need better DNA extraction/recovery
- Need further efforts with evidence interpretation and understanding of data being generated with high sensitivity techniques (move towards probabilistic methods)

We Need Continued Efforts to Improve DNA Interpretation (especially low-level DNA and mixtures)

Forensic Science International: Genetics 6 (2012) 677–678

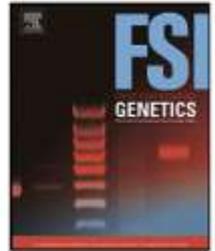


ELSEVIER

Contents lists available at SciVerse ScienceDirect

Forensic Science International: Genetics

journal homepage: www.elsevier.com/locate/fsig



Editorial

Focus issue—Analysis and biostatistical interpretation of complex and low template DNA samples

December 2012 – Forensic Science International: Genetics, volume 6, issue 6

Forensic Science International: Genetics 6 (2012) 679–688



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journal homepage: www.elsevier.com/locate/fsig



DNA commission of the International Society of Forensic Genetics:
Recommendations on the evaluation of STR typing results that may include drop-out and/or drop-in using probabilistic methods

P. Gill^{a,b,*}, L. Gusmão^c, H. Haned^d, W.R. Mayr^e, N. Morling^f, W. Parson^g, L. Prieto^h,
M. Prinzⁱ, H. Schneider^j, P.M. Schneider^k, B.S. Weir^l

Summary of DNA Mixture Interlaboratory Studies Conducted by NIST

Study	Year	# Labs	# Samples	Mixture Types
MSS 1	1997	22	11 stains	ss, 2p, 3p
MSS 2	1999	45	11 stains	ss, 2p, 3p
MSS 3	2000-01	74	7 extracts	ss, 2p, 3p
MIX05	2005	69	4 cases (.fsa)	only 2p
MIX13	2013	108	5 cases (.fsa)	2p, 3p, 4p

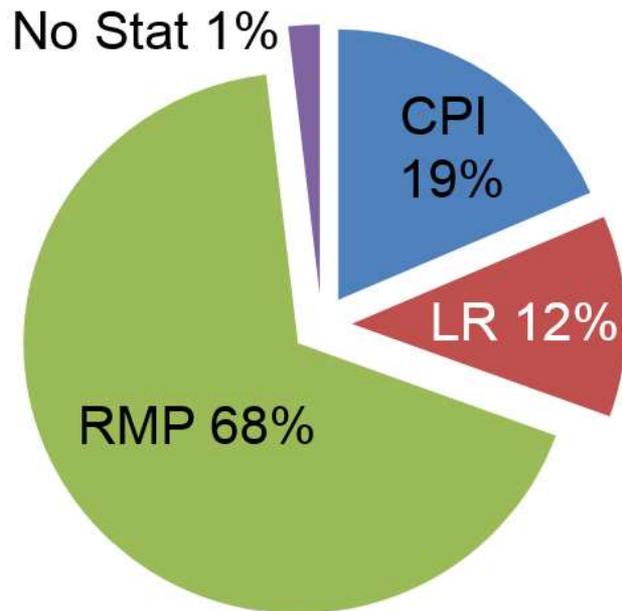
ss = single-source
2p = 2-person
3p = 3-person
4p = 4-person

- Other recent studies
 - UK Regulator
 - USACIL

Studies have revealed significant variations in approaches among and within forensic laboratories

MIX13 Study Case 1 Results

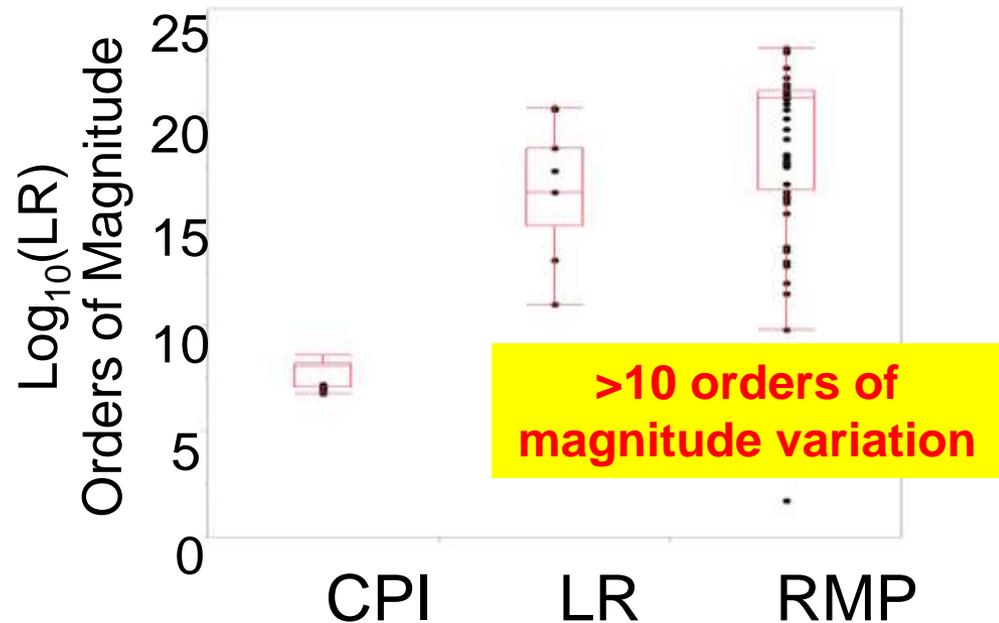
- **Scenario:** Mock sexual assault, **2 person 50:50 mixture**, all alleles above a stochastic threshold of 150 RFU
- **Purpose:** How many labs would consider the victim's profile and determine genotypes through deconvolution and report a modified random match probability statistic?



N = 108 laboratories

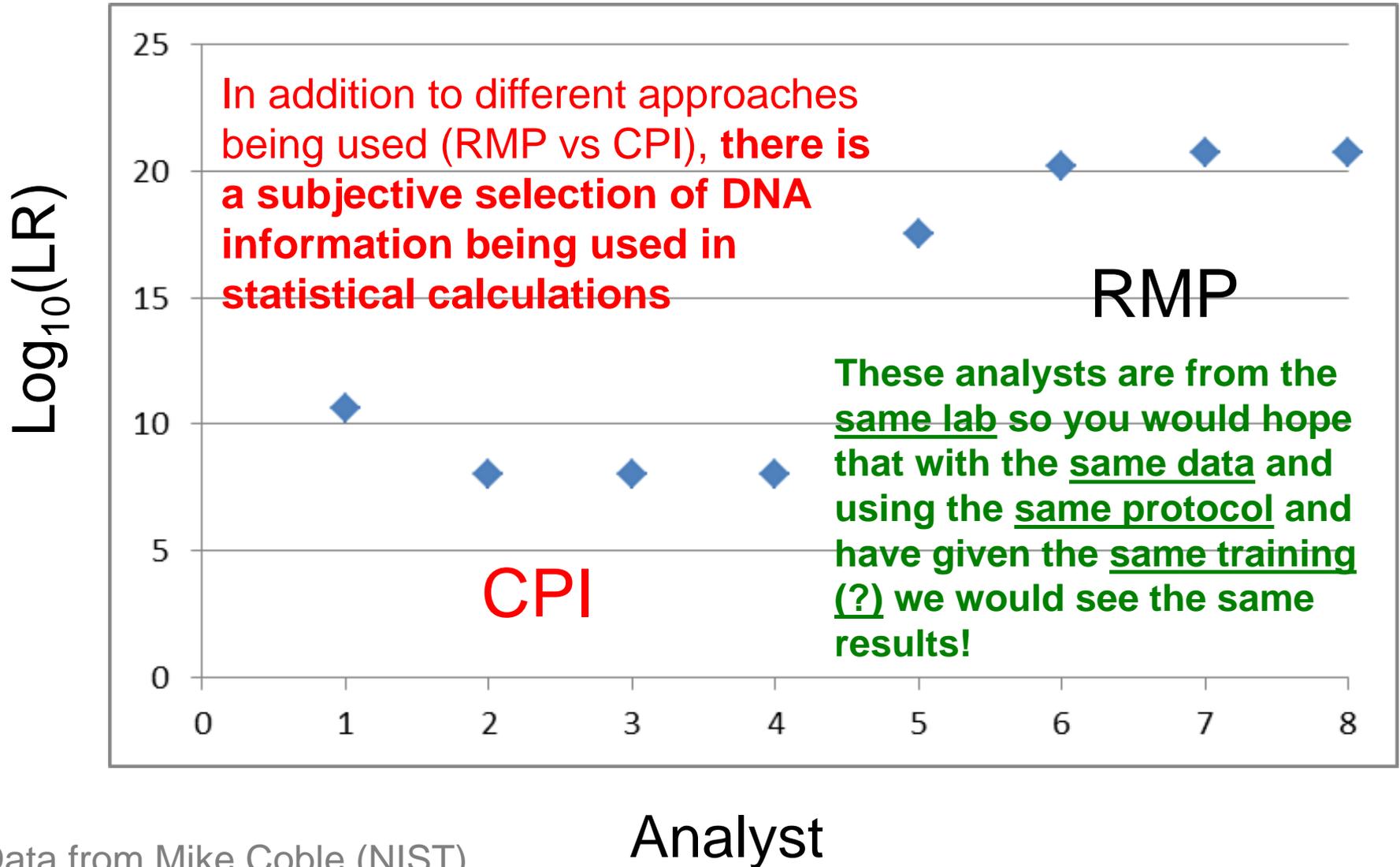
Data from Mike Coble (NIST)

Different Approaches Being Taken



MIX13 Study Case 1 Results

Intra-Laboratory Results (n = 8)



DNA Interpretation Training Workshops

melbourne

25th World Congress of the International
Society for Forensic Genetics
2 – 6 September 2013

September 2-3, 2013

Two days of basic and advanced workshops
on DNA evidence interpretation



Handouts and reference list available at

<http://www.cstl.nist.gov/strbase/training/ISFG2013workshops.htm>

The Workshop Instructors



**Mike Coble
(NIST)**

**Peter Gill
(U. Oslo)**

**Jo Bright
(ESR)**

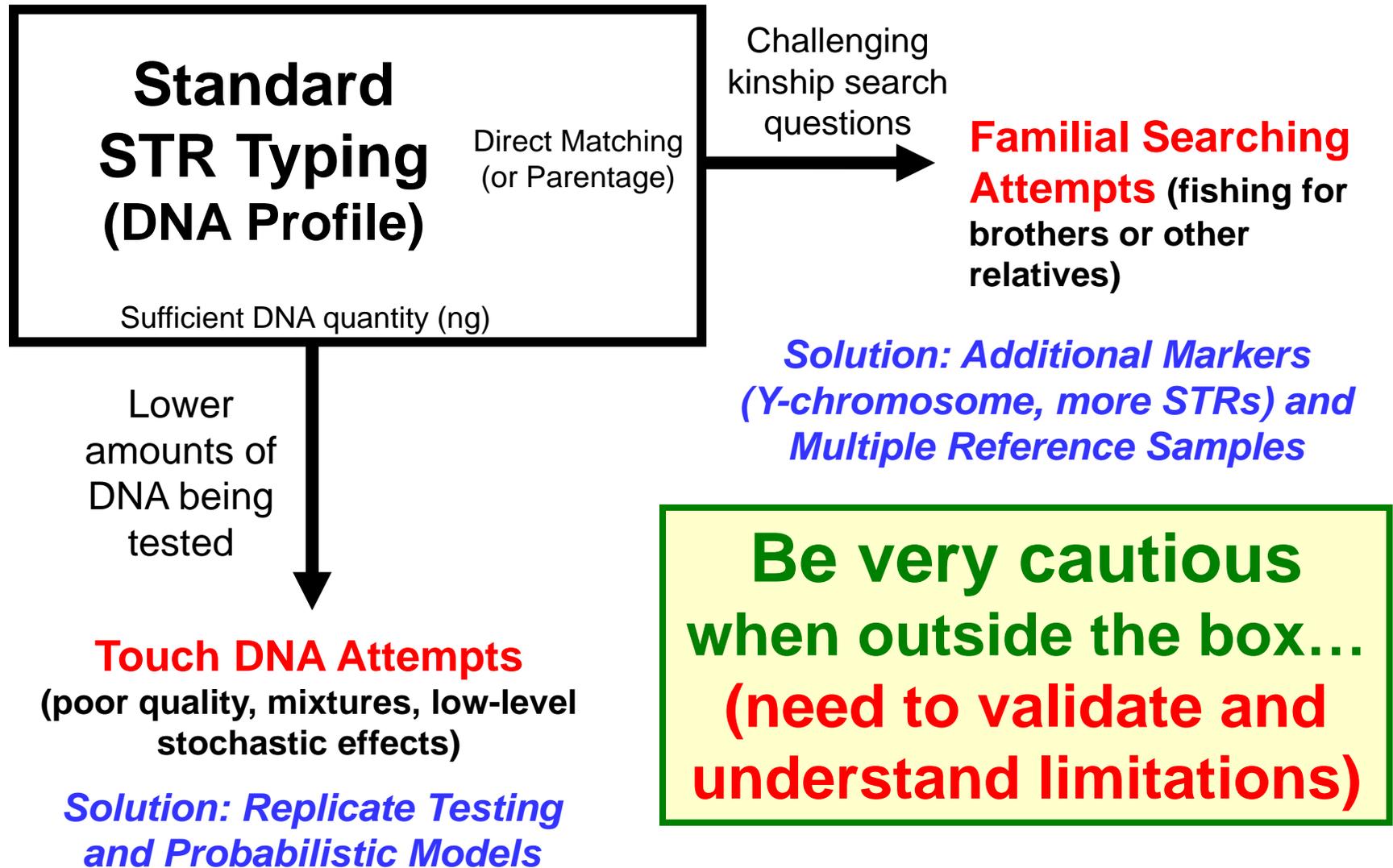
**John Buckleton
(ESR)**

**Duncan Taylor
(FSSA)**

**John Butler
(NIST)**

Going Beyond the Core Competencies of Forensic DNA Testing...

Core Competency



Some Thoughts on the Future...

- **PCR amplification**
 - Faster enzymes to enable rapid PCR
 - More robust enzymes and master mixes to overcome inhibition
- **Instrumentation**
 - More dye colors to aid higher levels of multiplexing
 - Rapid, integrated devices
 - Alternatives to capillary electrophoresis: ~~PLEX-ID~~ and NGS
- **Quantitative information**
 - qPCR and digital PCR
- **Marker systems**
 - Expanding sets of STR loci for growing DNA databases
 - Other marker systems: SNPs, InDels, X-STRs, RM Y-STRs
 - Body fluid identification with mRNA, miRNA, and DNA methylation
 - Phenotyping for external visible characteristics
 - Challenges with potential whole genome information
- **Data interpretation**
 - Probabilistic genotyping for low-level DNA and mixture interpretation
 - Probability of dropout

The Future of Forensic DNA

is Similar to the Olympic Motto of
“Swifter, Higher, Stronger”

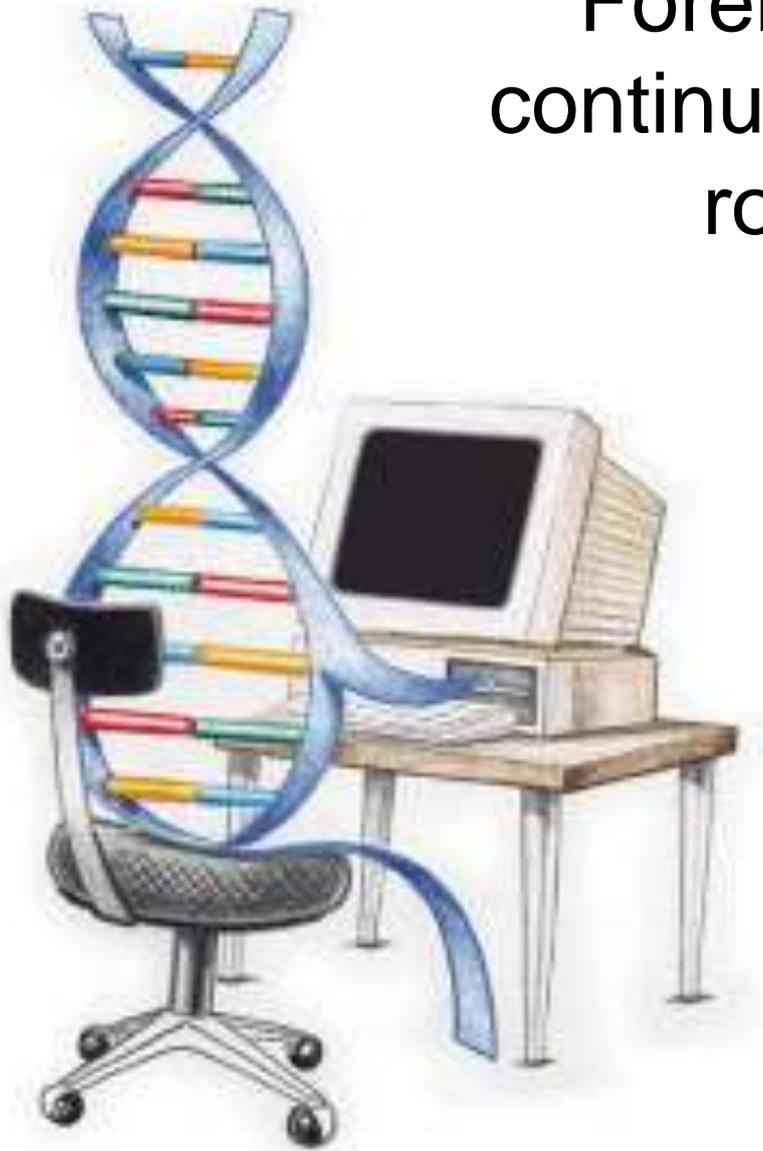


Resources

Training

Action

Forensic DNA testing will continue to play an increasing role in the future...



<http://www.manastungare.com/publications/genetic/dna.gif>

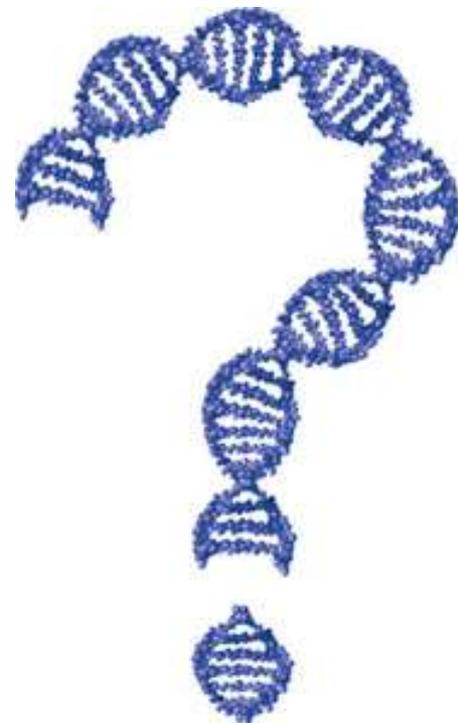
- Improved sensitivities lead to contamination concerns and DNA case relevance
- Costs will play a role
 - Competition
 - Centralization
 - Communication
- New technology adoption?
 - Political factors
 - Legal factors
 - Legacy data
 - Privacy concerns

National Commission on Forensic Science (NCFS):
www.justice.gov/ncfs

Organization of Scientific Area Committees (OSAC):
www.nist.gov/forensics/osac/index.cfm



www.nist.gov/forensics



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Mathematical Analogy to Forensic DNA

- **Forensic DNA testing** can be equated to different levels of **mathematics**:
 1. **Single-source samples** (e.g., reference samples) are like **basic arithmetic**
 2. **Two-person mixtures** (e.g., sexual assault evidence) are like **algebra**
 3. **Complex mixtures** (e.g., touch DNA) are like **calculus**
- Validation studies can be considered classroom instruction to help understand the topic and prepare for the casework “exams”
- Proficiency tests are like homework – a graded experience where feedback is received to prepare students for the casework exams when the true answers are not known to the test takers
- If homework is not challenging enough or if your classroom instruction is not to the level needed to be prepared, how can you hope to pass the test? Algebraic principles are necessary for calculus (just like two-person mixture principles apply to complex mixtures), but to truly solve calculus problems and complex mixtures a different level of thinking is required and more study is necessary. ***Would you want to go into a calculus final with only instruction in algebra and experience doing only basic math homework problems?***